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FIG. 1A

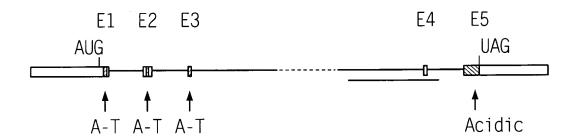


FIG. 1B





FIG. 2A

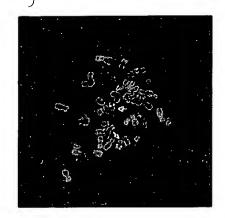


FIG. 2B



FIG. 2C

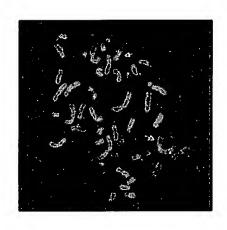


FIG. 2D

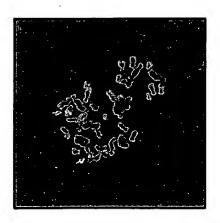


FIG. 2E

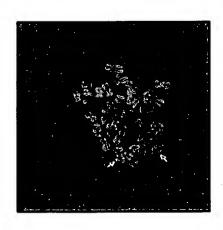


FIG. 2F



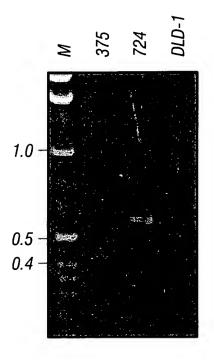


FIG. 3



HMGI-C AGA GGC AGA CCT AGG AAA TGG CCA CAA CAA GTC P Q Q V

fusion sites

t(3;12) AGA GGC AGA CCT AGG AAA TGG AAT ACT CTG GA N T L E

t(12;15) AGA GGC AGA CCT AGG AAA TGG CCC GGG CTC CAG R P R K W P G L Q

FIG. 4



RNA	ST90- 375	ST93- 724
Primer	375 724	724 375

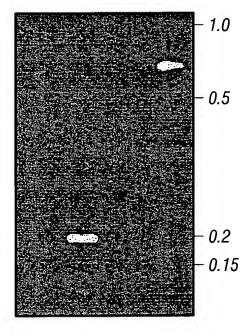


FIG. 5



```
Chr.3: 1 QCNVCSKPI.[x9]..KAYHPHCFTCVMCHRSL.[x11].LIHCIEDF Zyxin: 2 KCSVCKQTI.[x9]..NSYHPQCFTCVMCHTPL.[x11].QPHCVDDY Ap: 1 DCSGCGRQI.[x10].KRWHASCLKCYACRQPL.[x11].NIYCKNDY Lh2: 1 LCAGCGGKI.[x10].KQWHMRCLKCCECKLNL.[x11].SIYCKEDY Lin11: 1 ECAACAQPI.[x10].KCWHQSCLRCCDCRAPM.[x9]..LILCKTDF RBTN-1: 1 GCAGCNRKI.[x10].KYWHEDCLKCACCDCRL.[x12].LILCRRDY
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Chr.3: 2 RCSVCKEPI.[x15].SRFHVHCYRCEDCGGLL.[x13].HILCKTCN Zyxin: 3 RCSVCSEPI.[x16].KNFHMKCYKCEDCGRPL.[x14].HVLCMKCH RCSRCLASI.[x11].LVFHVNCFCCTVCH.PL.[x11].LIYCRTHY RCARCHLGI.[x11].LVYHNLCFTCTTCN.ML.[x11].LVYCRLHF Lin11: 2 RCAGCDGKL.[x11].KVFHIRCFQCSVCQRLL.[x12].RFVCQSDF RBTN-1: 2 NCAACSKLI.[x11].NVYHLDCFACQLCNQRF.[x12].IL.CQMDY
```

Consensus CxxC x16-21 HxxCxxCxxC x16-21 CxxH

FIG. 6A

<u>Chr. 15</u> <u>EEEE</u>H LNT<u>E</u>RSSAGGGWRGVQPLGS PTPG<u>ED</u>HRPIPSPASGFPSI

FIG. 6B



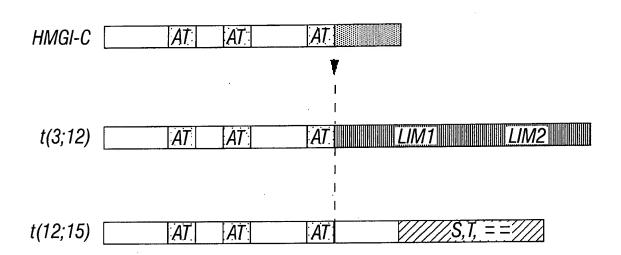
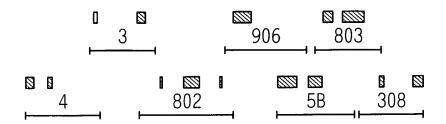


FIG. 7



PYGMY LOCUS



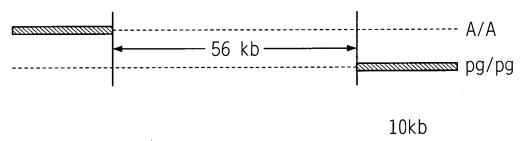


FIG. 8A

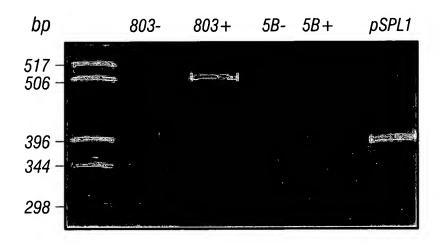


FIG. 8B



HIV-tat

AAGCAGCAGCAAGAGCCAACCTGTGAGCCCTCTCCTAAGAGACCCAGAGGAAGACCCAAA TCTATCAAAGCAGAGCCAACCTGTGAGCCCTCTCCTAAGAGACCCAGAGGAAGACCCAAA

GGCAGCAAAAACAAGAGCCCCTCTAAAGCAGCCCAGAAGAAAGCAGAGAGACCATTGGAGAA 61

HIV-tat

Mouse HMG I-C 803 121 AAACGGCCAAGAGGCAGACCTAGGAAATGGACCCACCTCCCA AAACGCCCAAGAGCCAGACCTAGGAAATGGCCACAACAAGTC

FIG. 8C

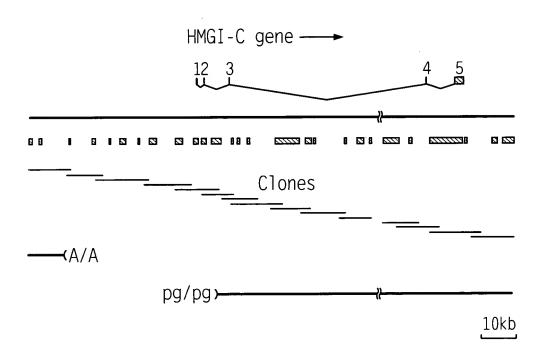


FIG. 8D



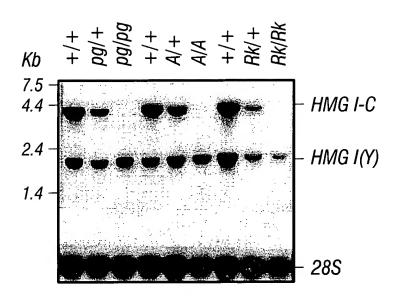


FIG. 9



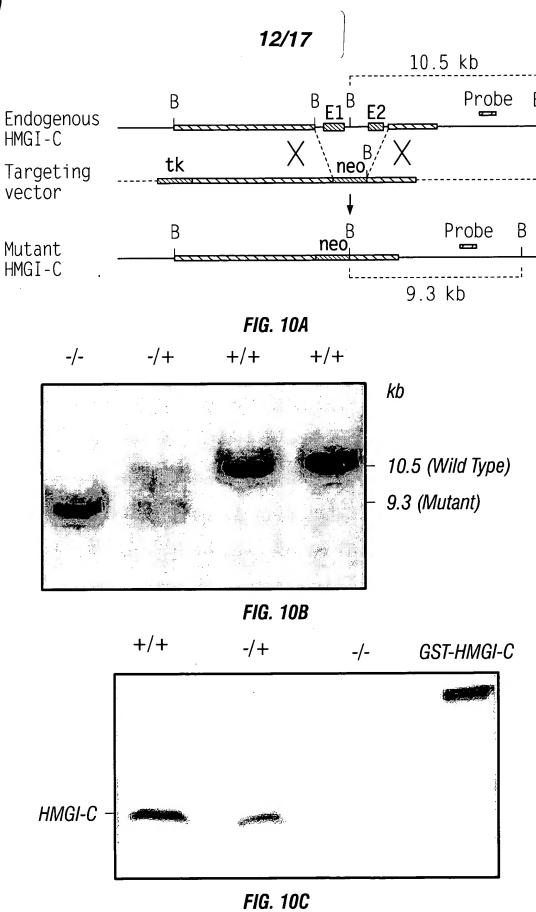






FIG. 11A



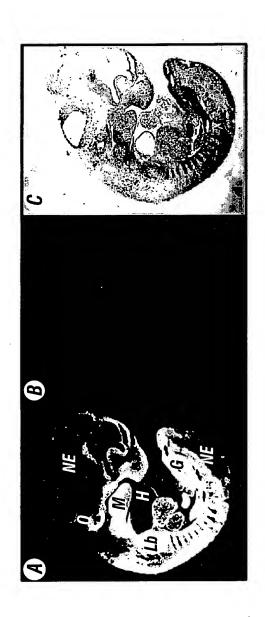


FIG. 11B

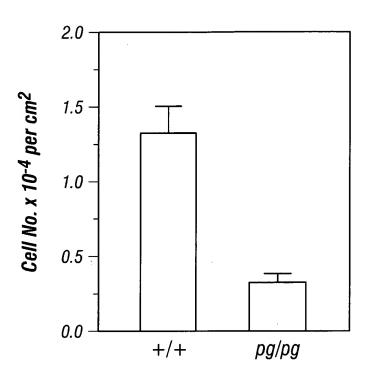


FIG. 11C

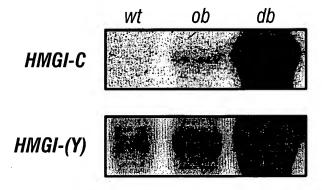


FIG. 12

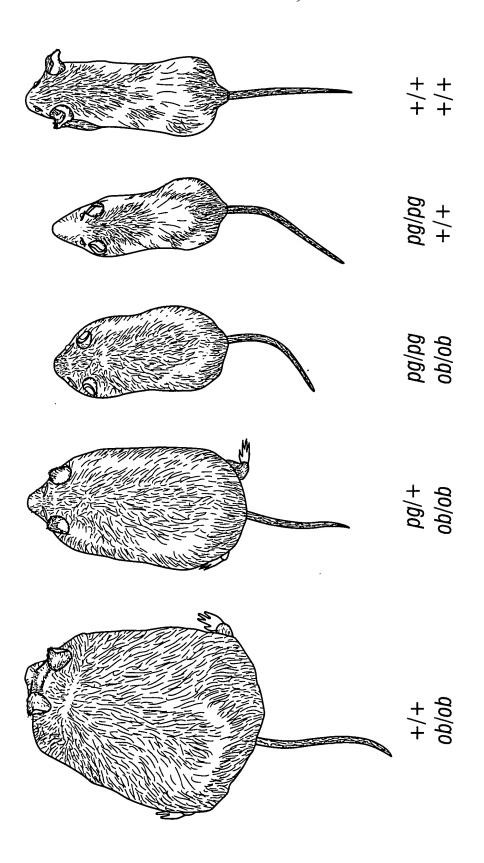


FIG. 13



